

Additional file 8: Table S4:

Motif Name	Query count	SD	P-value	Motif seq
Constitutive BDPs				
SORLIP2AT	500	5.146584	1.00e-7	GGGCC
SITEIIATCYTC	47	10.58654	0	TGGGCY
UP1ATMSD	81	9.883992	0	GGCCCAWWW
Seedling-specific BDPs				
TBF1HSF	17	2.625993	4.32e-3	GAAGAAGAA
Callus-specific BDPs				
ACGTABREMOTIFA2OSEM	41	3.206062	6.73e-4	ACGTGKC
BOXIIPCCHS	31	3.055217	1.12e-3	ACGTGGC